

## SEQUENCE LISTING

<110> Zinselmeier, Chris  
 Habben, Jeff  
 Tomes, Dwight

<120> Regulated Expression of Genes in Plant  
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96

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 Pro Trp Pro Ala Phe Leu Ala Ala Leu Ala Asp Gly Lys Leu Arg  
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144

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 Thr Asp Ser Asn Ala Thr Ala Ala Ser Thr Asp Phe Gly Asn Ile  
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192

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240

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288

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336

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384

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432

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| Met Phe Val Pro Arg Ser Arg Ile Ala Asp Phe Asp Arg Gly Val Phe |     |     |      |
| 405   | 410 | 415 |      |
| aag ggc atc ttg cag ggc acc gac atc gtc ggc ccg ctc atc gtc tac |     |     | 1296 |
| Lys Gly Ile Leu Gln Gly Thr Asp Ile Val Gly Pro Leu Ile Val Tyr |     |     |      |
| 420   | 425 | 430 |      |
| ccc ctc aac aaa tcc atg tgg gac gac ggc atg tcg gcg gcg acg ccg |     |     | 1344 |
| Pro Leu Asn Lys Ser Met Trp Asp Asp Gly Met Ser Ala Ala Thr Pro |     |     |      |
| 435   | 440 | 445 |      |
| tcg gag gac gtg ttc tac gcg gtg tcg ctc ttc tcg tcg gtg gcg     |     |     | 1392 |
| Ser Glu Asp Val Phe Tyr Ala Val Ser Leu Leu Phe Ser Ser Val Ala |     |     |      |
| 450   | 455 | 460 |      |
| ccc aac gac ctg gcg agg ctg cag gag cag aac agg agg atc ctg cgc |     |     | 1440 |
| Pro Asn Asp Leu Ala Arg Leu Gln Glu Gln Asn Arg Arg Ile Leu Arg |     |     |      |
| 465   | 470 | 475 | 480  |
| ttc tgc gac ctc gcc ggg atc cag tac aag acc tac ctg gcg cgg cac |     |     | 1488 |
| Phe Cys Asp Leu Ala Gly Ile Gln Tyr Lys Thr Tyr Leu Ala Arg His |     |     |      |
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| acg gac cgc agt gac tgg gtc cgc cac ttc ggc gcc gcc gag tgg aat |     |     | 1536 |
| Thr Asp Arg Ser Asp Trp Val Arg His Phe Gly Ala Ala Glu Trp Asn |     |     |      |
| 500   | 505 | 510 |      |
| cgc ttc gtg gag atg aag aac aag tac gac ccc aag agg ctg ctc tcc |     |     | 1584 |
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| ccc ggc cag gac atc ttc aac tga                                 |     |     | 1608 |
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| Pro Trp Pro Ala Phe Leu Ala Ala Leu Ala Leu Asp Gly Lys Leu Arg |     |     |      |
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| 50  | 55  | 60  |      |
| Thr Ser Ala Leu Pro Ala Ala Val Leu Tyr Pro Ser Ser Thr Gly Asp |     |     |      |
| 65  | 70  | 75  | 80   |
| Leu Val Ala Leu Leu Ser Ala Ala Asn Ser Thr Pro Gly Trp Pro Tyr |     |     |      |
| 85  | 90  | 95  |      |
| Thr Ile Ala Phe Arg Gly Arg Gly His Ser Leu Met Gly Gln Ala Phe |     |     |      |
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 His Gly Pro Gln Ile Ser Asn Val Leu Glu Met Asp Val Ile Thr Gly  
 195 200 205  
 His Gly Glu Met Val Thr Cys Ser Lys Gln Leu Asn Ala Asp Leu Phe  
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 Asp Ala Val Leu Gly Gly Leu Gly Gln Phe Gly Val Ile Thr Arg Ala  
 225 230 235 240  
 Arg Ile Ala Val Glu Pro Ala Pro Ala Arg Ala Arg Trp Val Arg Leu  
 245 250 255  
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 Ala Pro Arg Pro Gly Gly Gly Ala Ser Phe Gly Pro Met Ser Tyr  
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 Val Glu Gly Ser Val Phe Val Asn Gln Ser Leu Ala Thr Asp Leu Ala  
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 305 310 315 320  
 Ala Gly Glu Arg Asn Ala Thr Thr Val Tyr Ser Ile Glu Ala Thr Leu  
 325 330 335  
 Asn Tyr Asp Asn Ala Thr Ala Ala Ala Val Asp Gln Glu Leu Ala  
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 Ser Val Leu Gly Thr Leu Ser Tyr Val Glu Gly Phe Ala Phe Gln Arg  
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 Asp Val Ser Tyr Thr Ala Phe Leu Asp Arg Val His Gly Glu Glu Val  
 370 375 380  
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 385 390 395 400  
 Met Phe Val Pro Arg Ser Arg Ile Ala Asp Phe Asp Arg Gly Val Phe  
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 Lys Gly Ile Leu Gln Gly Thr Asp Ile Val Gly Pro Leu Ile Val Tyr  
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 Pro Leu Asn Lys Ser Met Trp Asp Asp Gly Met Ser Ala Ala Thr Pro  
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 Pro Asn Asp Leu Ala Arg Leu Gln Glu Gln Asn Arg Arg Ile Leu Arg  
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| <223> Promoter and terminator from Zea mays as found in Genbank Accession #S78780; gene from Agrobacterium tumefaciens as found in Molecular and General Genetics 216:388-394 (1989). |      |
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| catgtgtga   | gcatataaga  | aacccttagt  | atgtatttgt | atttgtaaaa | tacttctatc  | 5580 |
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<211> 2722

<212> DNA

<213> Artificial Sequence

<220>

<223> Promoter from *Hordeum vulgare*, Plant Journal 6:849-860 (1994); gene from *Agrobacterium tumefaciens*, Molecular and General Genetics 216:388-394 (1989); terminator from *Zea mays*, Genbank Accession #S78780.

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| tcgtgagaat  | aaccgtggcc  | taaaaataag | ccgatgagga  | taaataaaat  | gtggtggatc   | 120  |
| agtacttcaa  | gaggttact   | catcaagagg | atgccttcc   | gatgagctct  | agtagtacat   | 180  |
| cgAACCTCAC  | atacctccat  | tgtggtaaaa | tattttgtgc  | tcatttagtgc | atgggtaaat   | 240  |
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| taatttccat  | tccggggcaa  | aagcaaaaca | attttatttt  | acttttacca  | ctcttagctt   | 360  |
| tcacaatgt   | tcacaaatgc  | cactctagaa | attctgttta  | tgccacagaa  | tgtaaaaaaaaa | 420  |
| aacactca    | tatttgaagc  | caagggtttc | atggcatgga  | aatgtgacat  | aaagtaacgt   | 480  |
| tcgtgtataa  | gaaaaaattt  | tactcctcg  | aacaagagac  | ggaaacatca  | tgagacaatc   | 540  |
| gcgtttggaa  | ggctttgcat  | caccttgg   | tgatgcgc    | aatggagtc   | gtctgcttgc   | 600  |
| tagcttcgc   | ctaccggcca  | ctgagtccgg | gcggcaacta  | ccatggcga   | acgaccccagc  | 660  |
| tgacctctac  | cgaccggact  | tgaatgcgc  | accttcgtca  | gchgacgttgg | ccgcgtacgc   | 720  |
| tggcgacgtg  | ccccggcatg  | catggggca  | catggcgagc  | tcagaccgtg  | cgtggctggc   | 780  |
| tacaaatacg  | tacccgtga   | gtgccctagc | tagaaactta  | cacctgcaac  | tgcgagagcg   | 840  |
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| aacaacaaca  | aacaacaac   | aacattacaa | ttactattta  | caattacagt  | cgacggatca   | 960  |
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| ttcgtttagt  | aattttgggg  | aaagcttctg | ccacagttt   | ttttcgatg   | aacagtggccg  | 1080 |
| cagtggcgct  | gatcttgc    | gtatccctgc | aatctgggt   | aacttatgtc  | tttttatatcc  | 1140 |
| ttcaactacca | tgaaaagact  | agtaatctt  | ctcgatgtaa  | catcgcc     | cactgctatt   | 1200 |
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| tctatcttcc  | ctgttctta   | atgaaagacg | tcattttcat  | cagtatgatc  | taagaatgtt   | 1320 |
| gcaacttgca  | aggaggcg    | tctttcttgc | aatttaacta  | actcggttgc  | tggccctgtt   | 1380 |
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| aacgcggatt  | ttcg        | tattattcgc | aacgagtttgc | cagacggag   | gagcttcatc   | 1920 |
| agcgtggcc   | agaccagat   | taagcagat  | ttacggcc    | ctgcagg     | gtcttattatc  | 1980 |
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| catgcgcgtc            | gacaggaaca            | gaaattccct | ttggggcg   | 2220 |
| gaaggaccac            | catttcaat             | gtgagttat  | ccccggcg   | 2280 |
| tgtgctgttag           | tatagccgt             | ggctagctag | ctagttgagt | 2340 |
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| aatagggtta            | tacaataatt            | gatatatgtt | ttctgttat  | 2520 |
| tttagatata            | gacaaaaaaa            | aatctaagaa | ctaaaacaaa | 2580 |
| agtatataatt           | gggataatgt            | cgatgagatc | cctcgtata  | 2640 |
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<212> DNA

<213> Artificial Sequence

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<223> Promoter from Zea mays, U.S. patent application 09/377,648;  
gene from Agrobacterium tumefaciens, Molecular and General  
Genetics 216:388-394 (1989); terminator from Solanum  
tuberosum, Plant Cell 1(1):115-122 (1989).

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| taatttccat  | tcccgccaa   | aagcaaaaca    | atttatTTT   | acttttacca  | ctcttagctt   | 360  |
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| tcgtgtataa  | gaaaaaattt  | tactcctcgt    | aacaagagac  | ggaaacatca  | tgagacaatc   | 540  |
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| tgacctctac  | cgaccggact  | tgaatgcgt     | accttcgtca  | gcgacgttgg  | cccggtacgc   | 720  |
| tggcgacgtg  | ccccggcatg  | catggcggca    | catggcgagc  | tcagaccgtg  | cgtggcttgc   | 780  |
| tacaaataacg | tacccctgt   | gtgccctagc    | tagaaactt   | cacctgcac   | tgcgagagcg   | 840  |
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<223> Synthesized based on sequence from Agrobacterium tumefaciens

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<223> Promoter from Zea mays, Genbank Accession #L22344;  
Gene from Agrobacterium tumefaciens, Molecular and  
General Genetics 216:388-394 (1989); terminator from  
Zea mays, Genbank Accession #L22345.

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